

Figure S1. Neighbor-joining phylogenetic tree of 23 *Pestalotiopsis* spp. isolates reconstructed from EF1- α sequences. Only bootstrap values > 50% calculated from 1,000 replications are shown. Serbian sequences are shown in bold. Accession numbers of the isolates retrieved from GenBank are in parentheses.

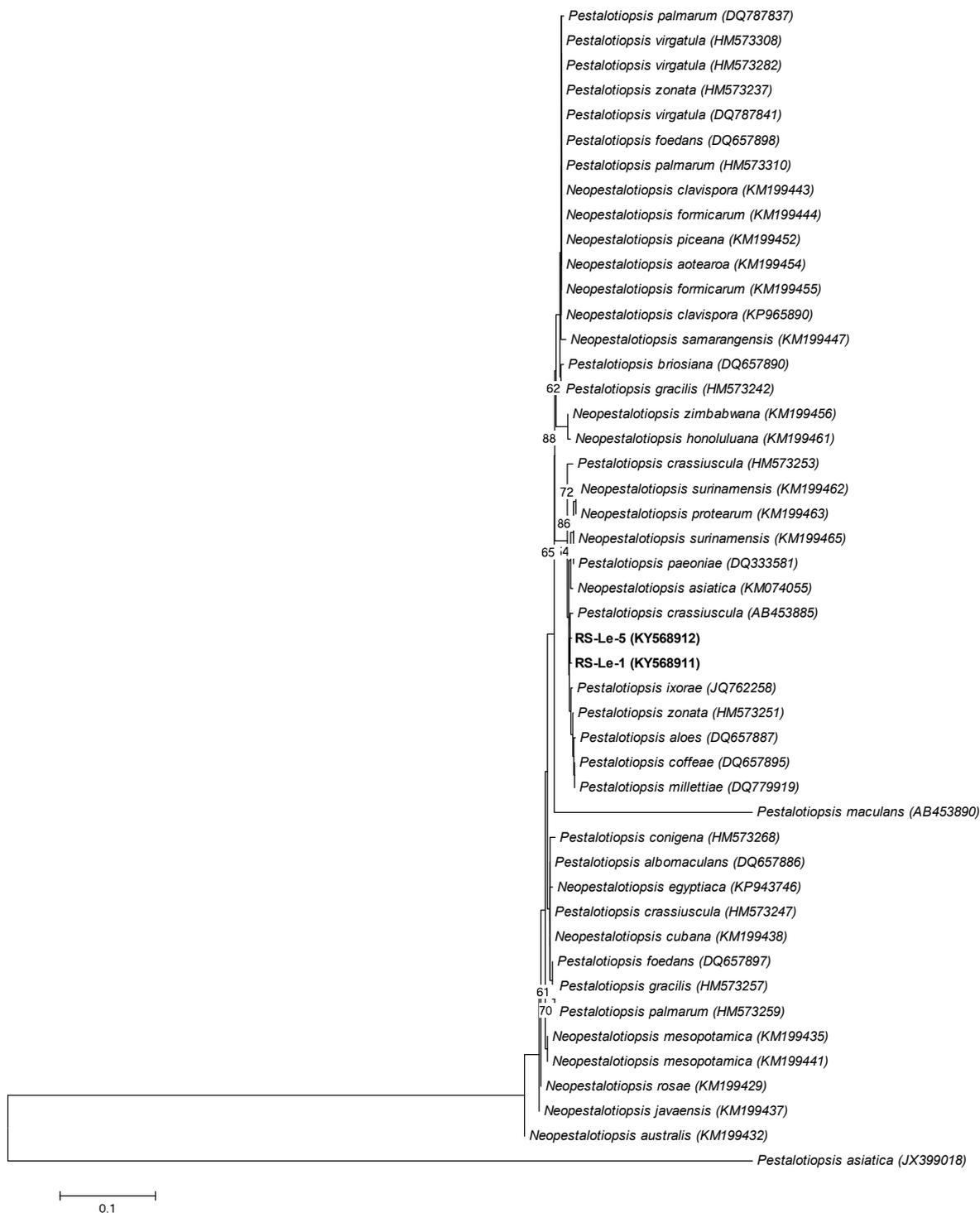


Figure S2. Neighbor-joining phylogenetic tree of 47 *Pestalotiopsis* spp. isolates reconstructed from TUB sequences. Only bootstrap values > 50% calculated from 1,000 replications are shown. Serbian sequences are shown in bold. Accession numbers of the isolates retrieved from GenBank are in parentheses.